

Table S1. CFU assay of WBM and splenocytes from WT or KO mice

Sources	Genotype	GEMM	GM	G	M	E	U	Total
BM	Tsc1 ^{fl/fl}	29.2 ± 5.6	229.2 ± 56.3	37.5 ± 6.4	50 ± 12.8	4.2 ± 1.3	0 ± 0	350 ± 45.3
	Tsc1 ^{fl/fl} ; Cre ⁺	116.7 ± 13.7**	179.2 ± 41.2	29.2 ± 11.2	150 ± 35.1**	8.3 ± 3.6	16.7 ± 8.6	500 ± 86.3*
Spleen	Tsc1 ^{fl/fl}	0 ± 0	5.8 ± 3.2	0 ± 0	5.8 ± 3.2	0 ± 0	0 ± 0	11.7 ± 6.5
	Tsc1 ^{fl/fl} ; Cre ⁺	5 ± 2.3**	24.2 ± 11.5	0.8 ± 0.6	18.3 ± 6.1*	0 ± 0	1.7 ± 1.2	50 ± 12.4**

Cells were harvested 10 d after plpC treatments and cultured in MethoCult GFM3434 medium for 12 d. The colonies were counted under a microscope based on their morphology. The numbers of colonies presented are mean and SEs of four independent experiments, normalized for every 100,000 cells. *, P < 0.05; **, P < 0.01. E, erythroid; G, granulocyte; GEMM, granulocyte erythroid macrophage and megakaryocyte; GM, granulocyte and macrophage; M, macrophage; U, unidentified.